

"The expert in anything was once a beginner"

CSIR NET – Life Science

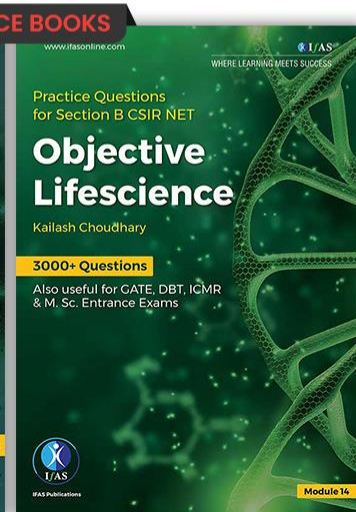
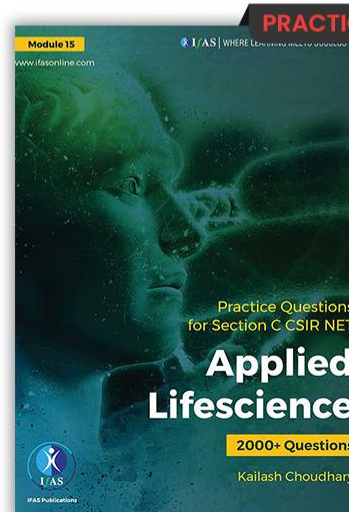
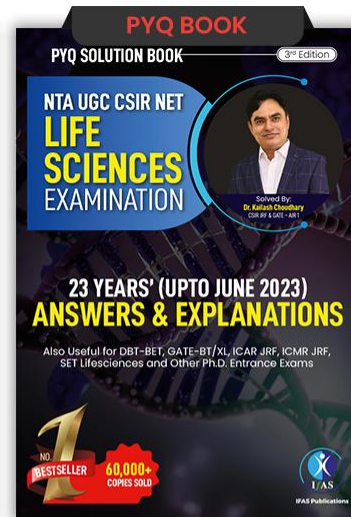
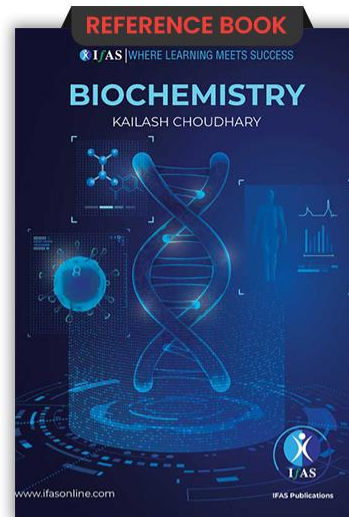
Unit 1: Biochemistry

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ETC and Oxidative Phosphorylation



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Points to be covered in this Lecture



Biological Oxidation



Redox Potential



Electron Transport Chain



Energy Transducer – ATP synthase



Oxidative Phosphorylation

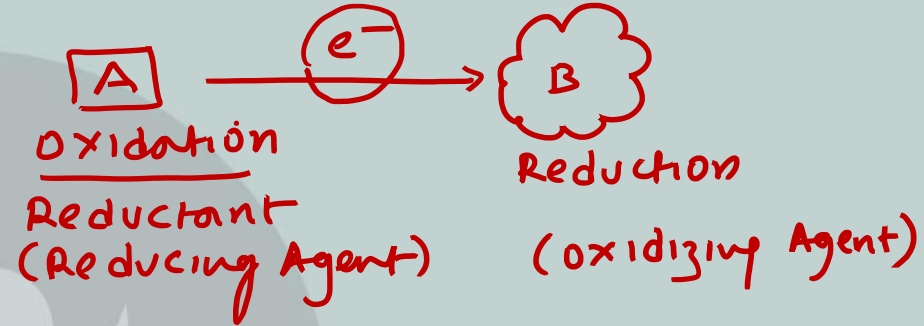
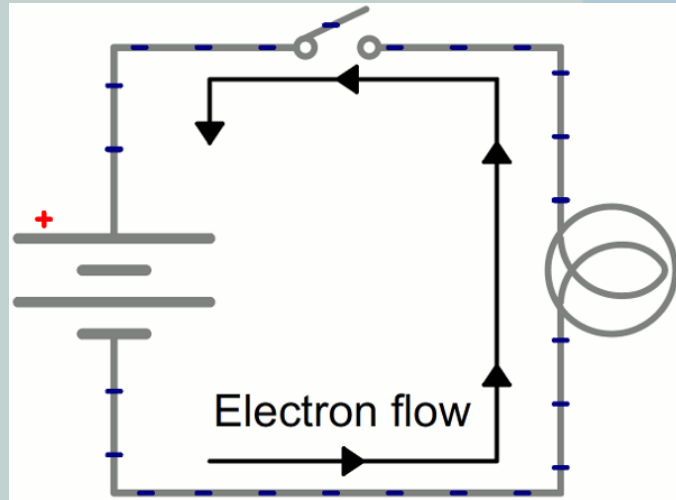


Shuttles



BIOLOGICAL OXIDATION

- ✓ Involves the loss of electrons
- ✓ Gain of oxygen or the loss of hydrogen.

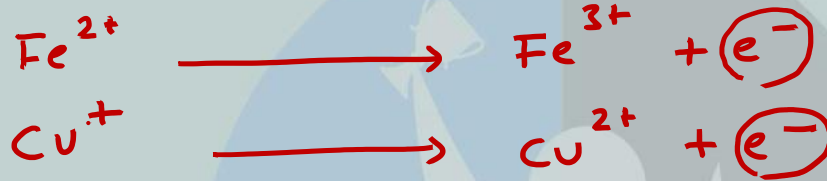


✓ The Flow of Electrons (emf) Can Do Biological Work

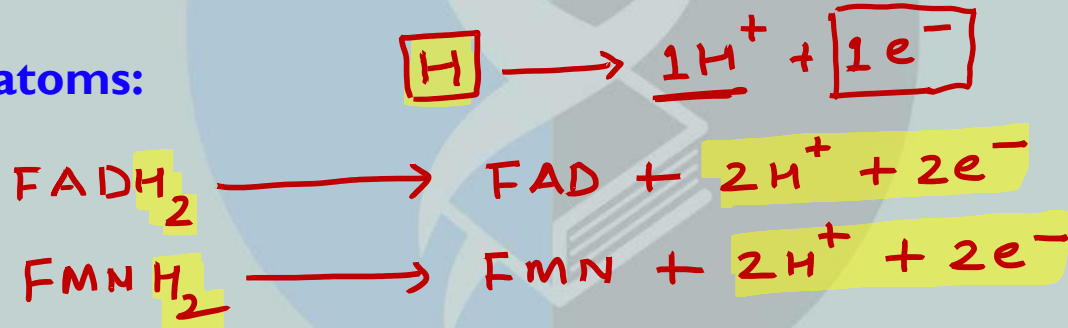


Four types of electron transfer occur in cells

1. Directly as electrons:



2. As hydrogen atoms:

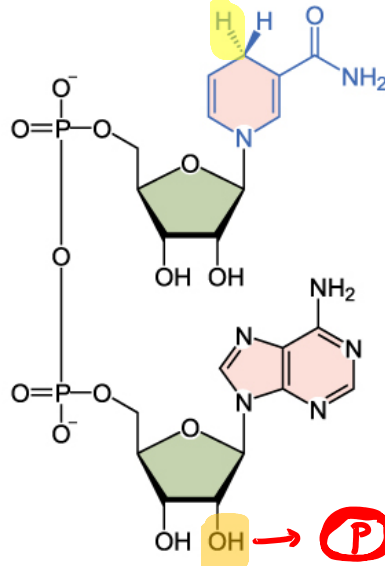
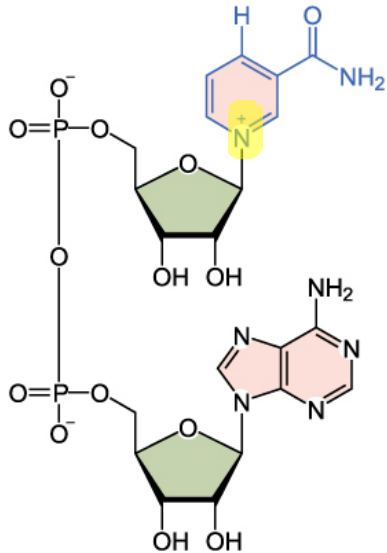




3. As a hydride ion (:H^-) which has two electrons:

NAD⁺

NADH





4. Through direct combination with oxygen:



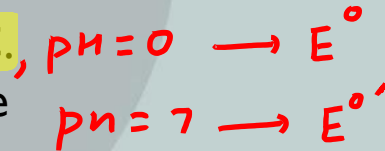
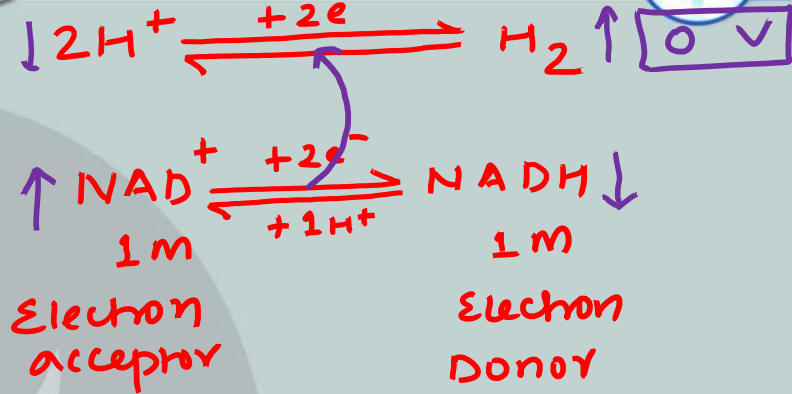


Reduction potential (measure in volts)

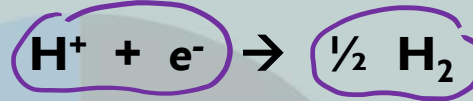
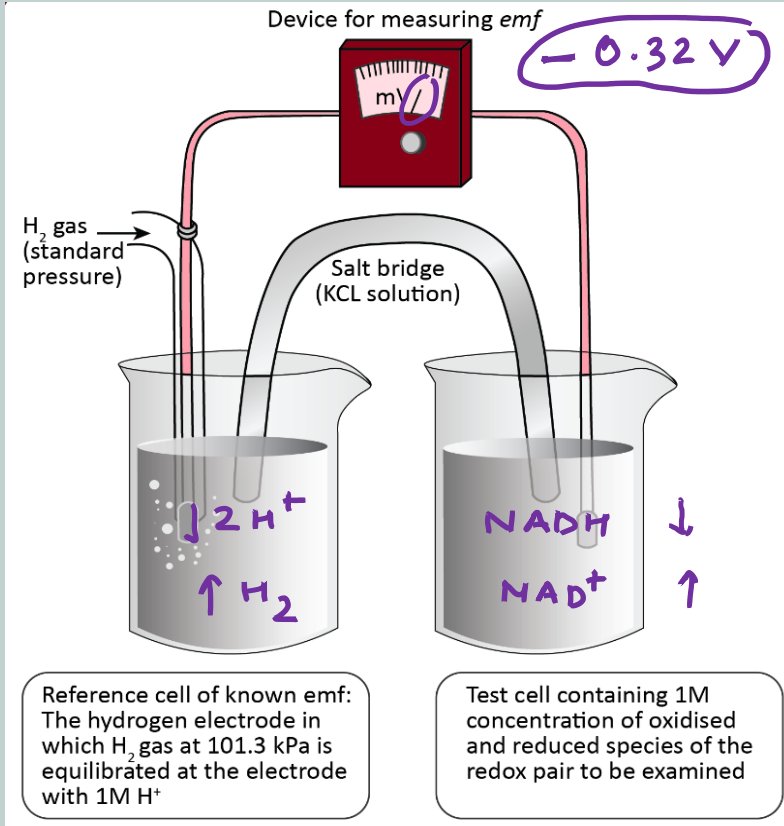
- Measure affinity for electrons
- Tendency of molecule to gain electrons

The standard reduction potential, E^0

- Reduction potential of a substance under standard conditions
- 1 M concentration, 1 atmospheric pressure, and 25°C .
- Measured relative to the standard hydrogen electrode (0 volts)



Standard reference



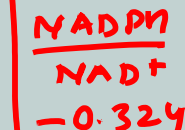
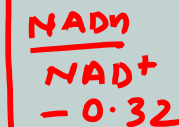
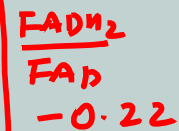
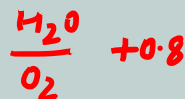
Test cell containing 1M (Oxidised/Reduced species)





Half-reaction	E'° (V)
$\frac{1}{2}\text{O}_2 + 2\text{H}^+ + 2e^- \longrightarrow \text{H}_2\text{O}$	0.816
$\text{Fe}^{3+} + e^- \longrightarrow \text{Fe}^{2+}$	0.771
$\text{NO}_3^- + 2\text{H}^+ + 2e^- \longrightarrow \text{NO}_2^- + \text{H}_2\text{O}$	0.421
Cytochrome <i>f</i> (Fe^{3+}) + $e^- \longrightarrow$ cytochrome <i>f</i> (Fe^{2+})	0.365
$\text{Fe}(\text{CN})_6^{3-}$ (ferricyanide) + $e^- \longrightarrow \text{Fe}(\text{CN})_6^{4-}$	0.36
Cytochrome <i>a</i> ₃ (Fe^{3+}) + $e^- \longrightarrow$ cytochrome <i>a</i> ₃ (Fe^{2+})	0.35
$\text{O}_2 + 2\text{H}^+ + 2e^- \longrightarrow \text{H}_2\text{O}_2$	0.295
Cytochrome <i>a</i> (Fe^{3+}) + $e^- \longrightarrow$ cytochrome <i>a</i> (Fe^{2+})	0.29
Cytochrome <i>c</i> (Fe^{3+}) + $e^- \longrightarrow$ cytochrome <i>c</i> (Fe^{2+})	0.254
Cytochrome <i>c</i> ₁ (Fe^{3+}) + $e^- \longrightarrow$ cytochrome <i>c</i> ₁ (Fe^{2+})	0.22
Cytochrome <i>b</i> (Fe^{3+}) + $e^- \longrightarrow$ cytochrome <i>b</i> (Fe^{2+})	0.077

Ubiquinone + $2\text{H}^+ + 2e^- \longrightarrow$ ubiquinol + H_2	0.045
Fumarate ²⁻ + $2\text{H}^+ + 2e^- \longrightarrow$ succinate ²⁻	0.031
$2\text{H}^+ + 2e^- \longrightarrow \text{H}_2$ (at standard conditions, pH 0)	0.000
Crotonyl-CoA + $2\text{H}^+ + 2e^- \longrightarrow$ butyryl-CoA	-0.015
Oxaloacetate ²⁻ + $2\text{H}^+ + 2e^- \longrightarrow$ malate ²⁻	-0.166
Pyruvate ⁻ + $2\text{H}^+ + 2e^- \longrightarrow$ lactate ⁻	-0.185
Acetaldehyde + $2\text{H}^+ + 2e^- \longrightarrow$ ethanol	-0.197
FAD + $2\text{H}^+ + 2e^- \longrightarrow$ FADH ₂	-0.219*
Glutathione + $2\text{H}^+ + 2e^- \longrightarrow$ 2 reduced glutathione	-0.23
$\text{S} + 2\text{H}^+ + 2e^- \longrightarrow \text{H}_2\text{S}$	-0.243
Lipoic acid + $2\text{H}^+ + 2e^- \longrightarrow$ dihydrolipoic acid	-0.29
$\text{NAD}^+ + \text{H}^+ + 2e^- \longrightarrow \text{NADH}$	-0.320
$\text{NADP}^+ + \text{H}^+ + 2e^- \longrightarrow \text{NADPH}$	-0.324
Acetoacetate + $2\text{H}^+ + 2e^- \longrightarrow$ β -hydroxybutyrate	-0.346
α -Ketoglutarate + $\text{CO}_2 + 2\text{H}^+ + 2e^- \longrightarrow$ isocitrate	-0.38
$2\text{H}^+ + 2e^- \longrightarrow \text{H}_2$ (at pH 7)	-0.414
Ferredoxin (Fe^{3+}) + $e^- \longrightarrow$ ferredoxin (Fe^{2+})	-0.432





Electron flow will be low redox potential to high redox potential
Standard reduction potentials can be used to calculate the free-energy change

	A		B
E°	-0.32	→	-0.22
E°	-0.3	→	0
E°	-0.3	→	+0.2
E°	+0.2	→	+0.3
	Low		High

$$\Delta E^\circ (E_B^\circ - E_A^\circ)$$

$$(-0.22) - (-0.32) = +0.1$$

$$(0) - (-0.3) = +0.3$$

$$(0.2) - (-0.3) = +0.5$$

$$(0.3) - (0.2) = +0.1$$

Spontaneous = $\Delta E^\circ = \text{positive change}$

$$\Delta G^\circ = -n F \Delta E^\circ$$

$$= -n \times 23 \times \Delta E^\circ (\text{Kcal})$$

$$= -n \times 96.5 \times \Delta E^\circ (\text{KJ/mole})$$

$n = \text{number of electron}$

$\text{Fe}^{2+} \text{ or } \text{Cu}^+ = 1 e^-$

$\text{FAD}, \text{NAD}^+, \text{NADP}, \text{UQ} = 2 e^-$

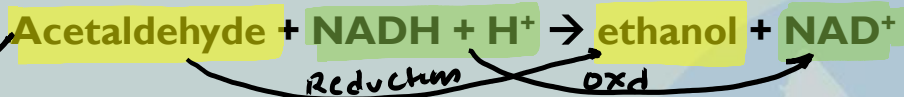
$\Delta G^\circ = \text{negative}$

$F = \text{Faraday constant } (96.5 \text{ kJ/V.mol or } 23 \text{ kcal/V.mol})$

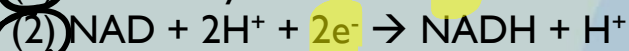
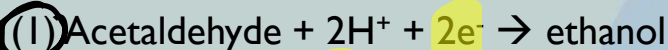


Consider the reaction in which acetaldehyde is reduced by the biological electron carrier NADH:

Electron Acceptor Electron donor



The relevant half-reactions and their E° values are:



Calculate ΔG° (Given $F = 96.5 \text{ kJ/V.mol}$)?

$$\begin{aligned} \Delta G^{\circ} &= -n F \Delta E^{\circ} \\ &= -2 \times 96.5 \times \Delta E^{\circ} \\ &= -2 \times 96.5 \times 0.1 \\ &= -193 \times 0.1 \end{aligned}$$

$$\Delta G^{\circ} = -19.3 \text{ kJ/mole}$$

standard condition

$$E^{\circ} = -0.2 \text{ V}$$

$$E^{\circ} = -0.3 \text{ V}$$

$$\begin{aligned} \Delta E^{\circ} &= (E^{\circ}_{\text{Acceptor}}) - (E^{\circ}_{\text{Donor}}) \\ &= (-0.2 \text{ V}) - (-0.3) \\ &= -0.2 + 0.3 \\ &= +0.1 \end{aligned}$$



Nerst Equation:

$$E = E^0 + \frac{0.26}{n} \ln \frac{(\text{Electron Acceptor})}{(\text{Electron Donor})}$$

- Cellular condition
- Physiological condition

$$\Delta G = -n F \Delta E$$

$$\Delta E = E_{\text{Acceptor}} - E_{\text{Donor}}$$



Electron Transport Chain

Location:

- Inner membrane of mitochondria
- Plasma membrane of bacteria

Components:

Four immobile complexes → I, II, III, IV

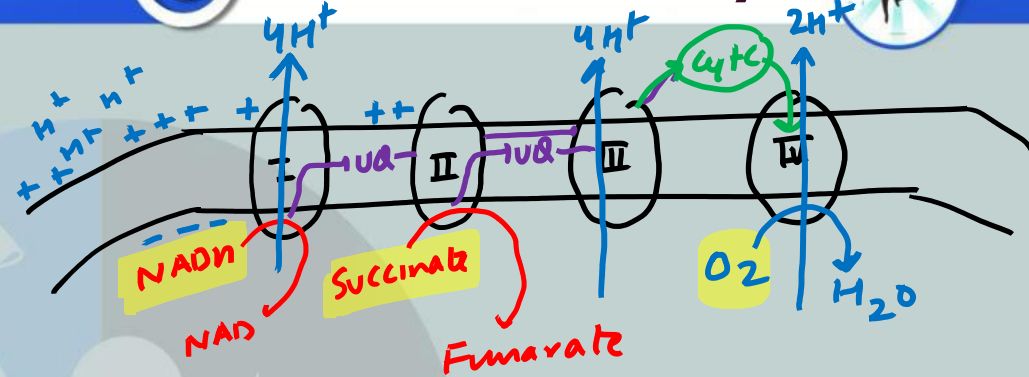
Ubiquinone (lipid soluble) and Cytochrome C (protein) are mobile

Functions

Oxidize **NADH** or **Succinate**

Transfer electron to oxygen

✓ Generate electrochemical gradient

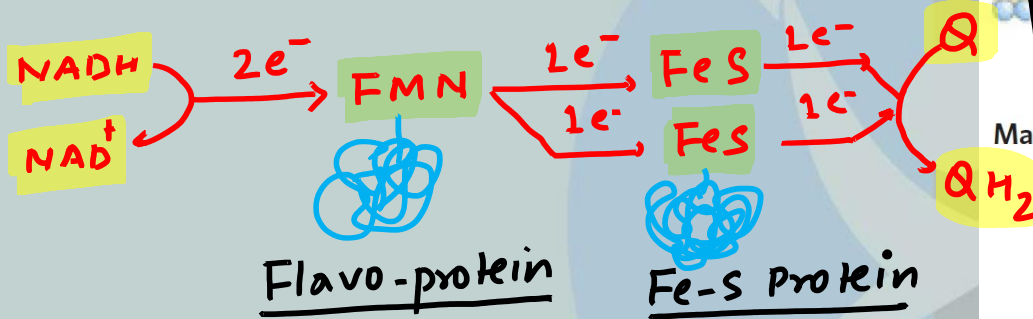


Electrical Gradient:
outside will be \oplus

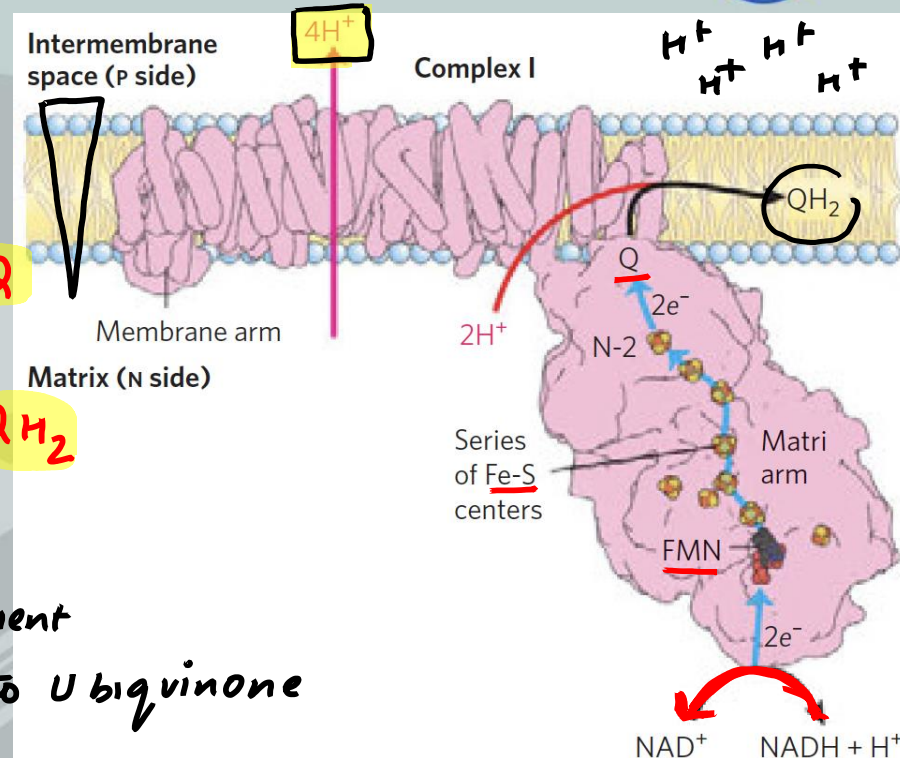
Chemical Gradient:
outside more H^+ conc^v

NADH-Q reductase (Complex I): NADH Dehydrogenase complex

- One molecule of flavin mononucleotide
- Eight or nine iron-sulfur clusters

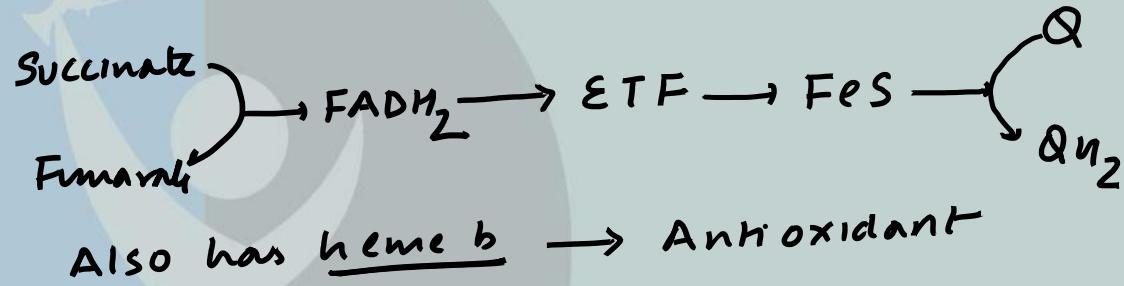
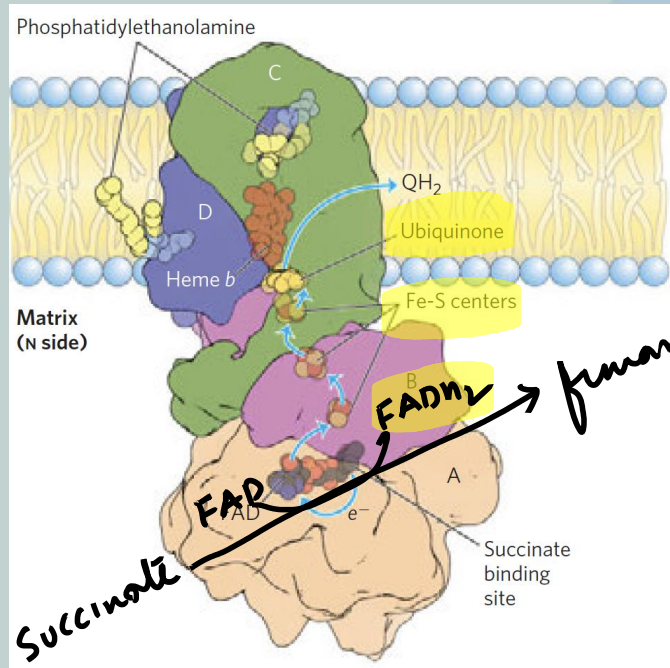


- 4H^+ are pumped against gradient during transfer $2e^-$ from NADH to Ubiquinone



Complex II: Succinate dehydrogenase complex

✓ Succinate to FAD to electron-transferring flavoprotein (ETF) ~~→~~ → Fes → Quinone
 Near **heme b** or the **quinone-binding site**: **Anti-oxidant** → **Prevents ROS**

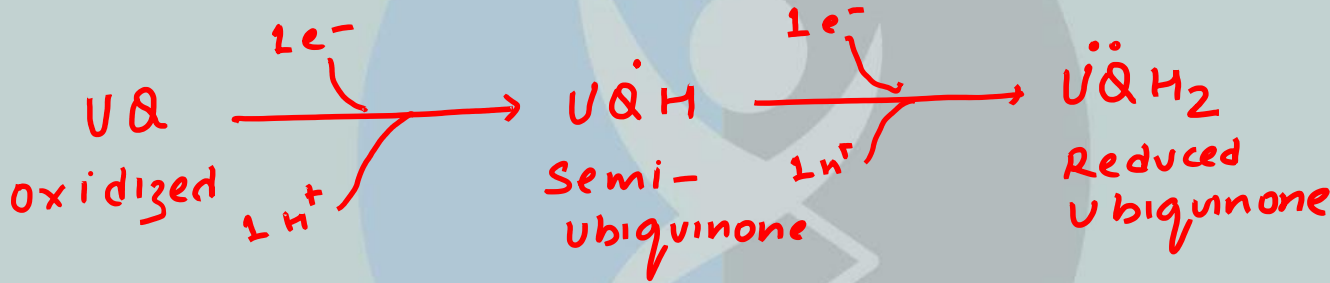




Ubi-Quinone

Lipid Soluble

- Ubiquinone (CoQ_{10}): Ten C5 isoprenoid units (50 carbon terpene — polyterpene)
- Transfer electron between complex I and III or II and III
- Protects cells from oxidative damage by neutralizing free radicals



Q-cytochrome c reductase (Complex III): Cyt b-c₁ complex

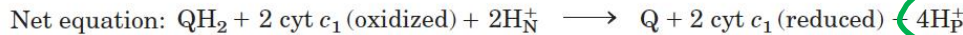
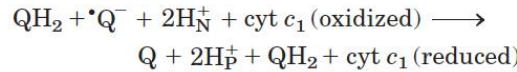
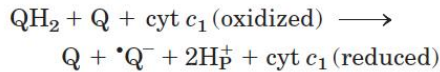
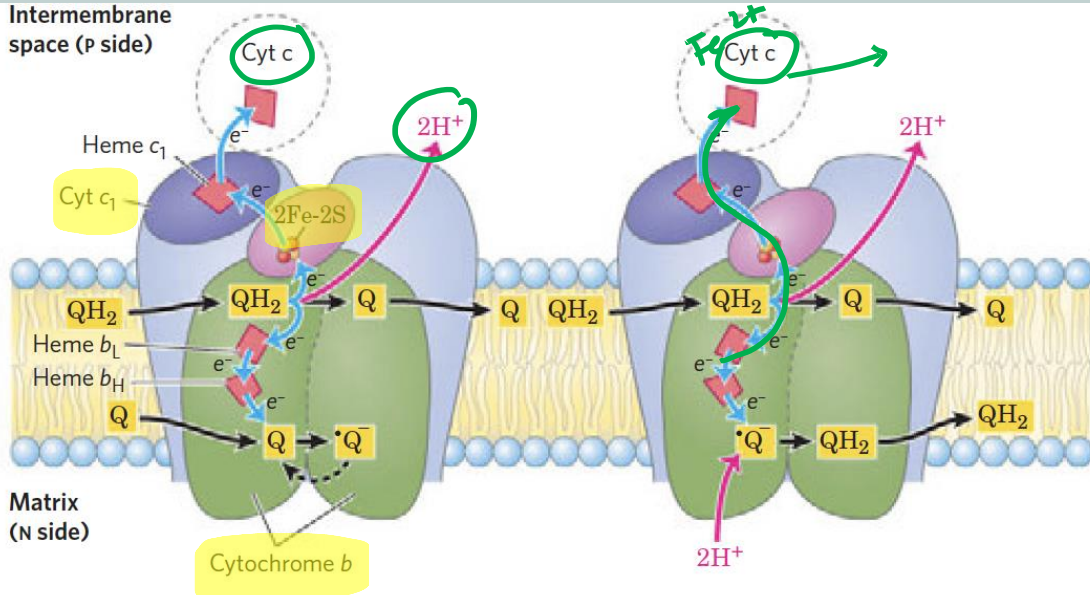
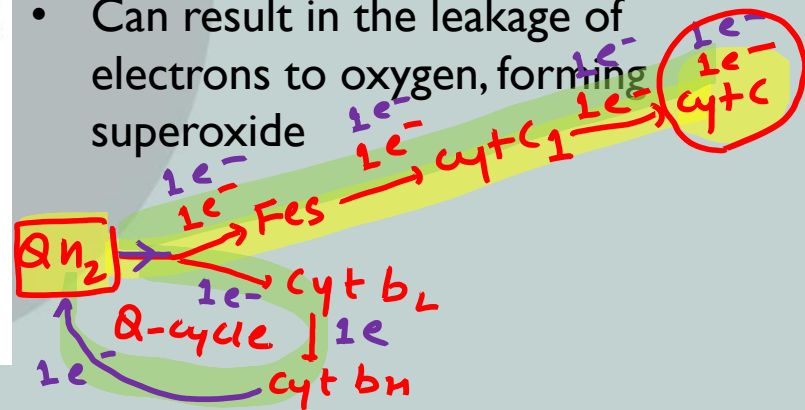
Haeme protein

- Consists of **cytochrome b**, **cytochrome c₁**, and the **Rieske iron-sulfur protein**.

- Pumping protons across the inner mitochondrial membrane

Q cycle = 4 H⁺ are pumped

- Can result in the leakage of electrons to oxygen, forming superoxide

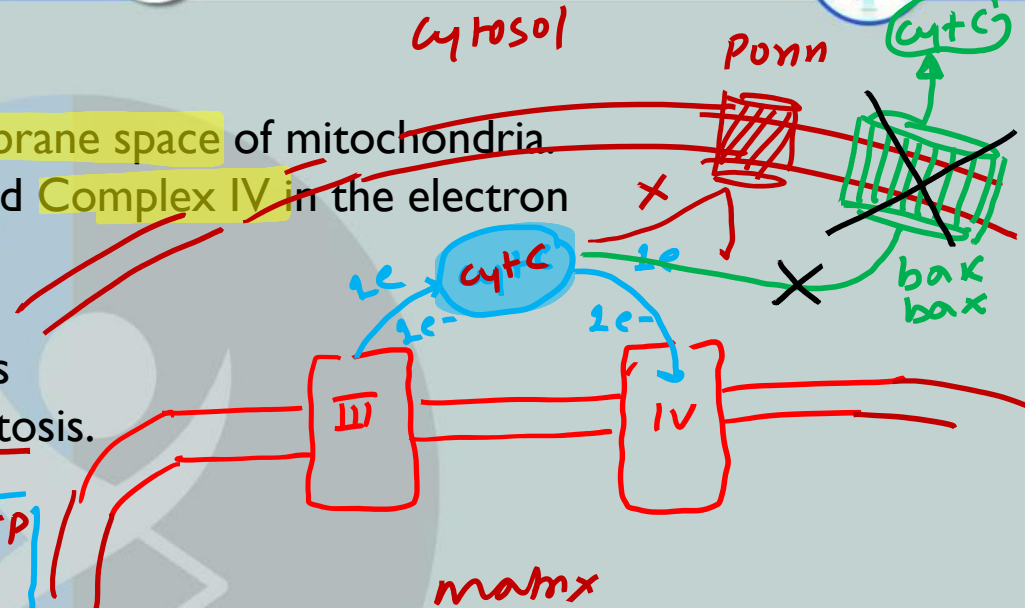




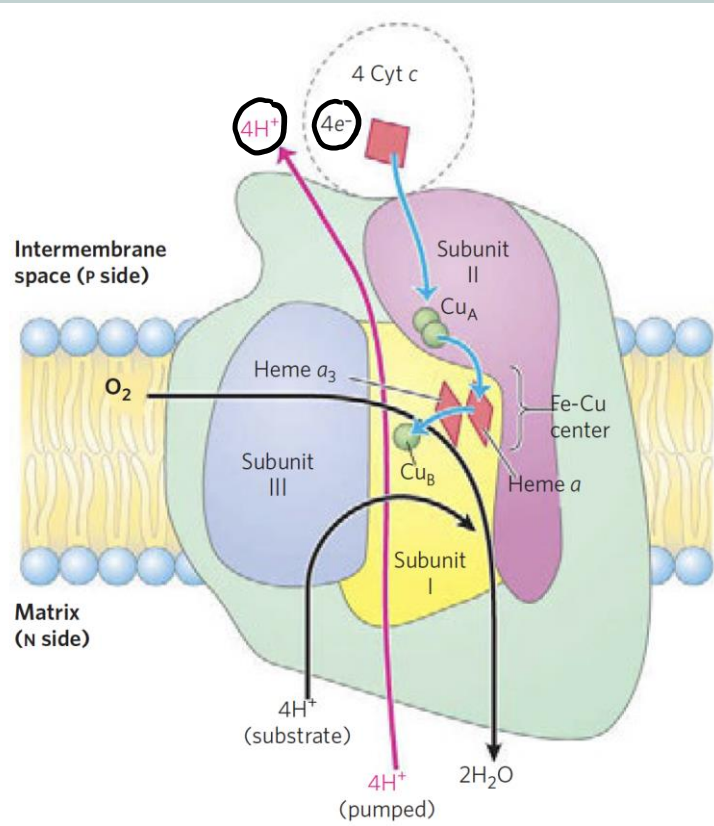
Cytochrome c

- Soluble protein located in the intermembrane space of mitochondria.
- Electron shuttle between Complex III and Complex IV in the electron transport chain.
- It contains a heme group
- Highly conserved across different species
- Involved in the intrinsic pathway of apoptosis.

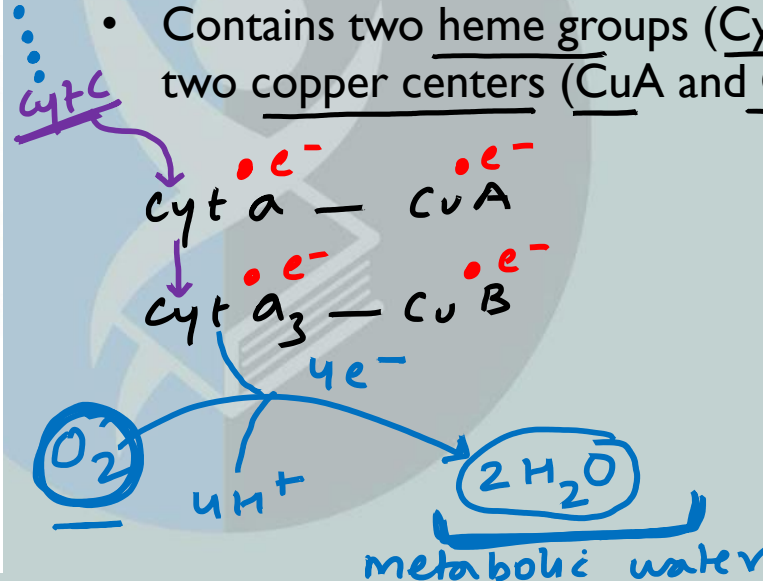
cytosol → $\text{cytc} + \text{Apaf-1} + \text{ATP} + \text{procaspase-9}$
Apoptosome complex



Cytochrome c oxidase (Complex IV):



- Transfer of electrons from cytochrome c to molecular oxygen
- Translocates protons across the inner mitochondrial membrane $(2e^-) \rightarrow (2H^+) \quad 4e^- \rightarrow 4H^+$
- Contains two heme groups (Cyt a and a3) and two copper centers (CuA and CuB).





Electron transport chain inhibitors:

Rotenone
Amytal
Barbiturates
Piercidin

⊥[⊖]
Complex-I

Malonate
Carboxin

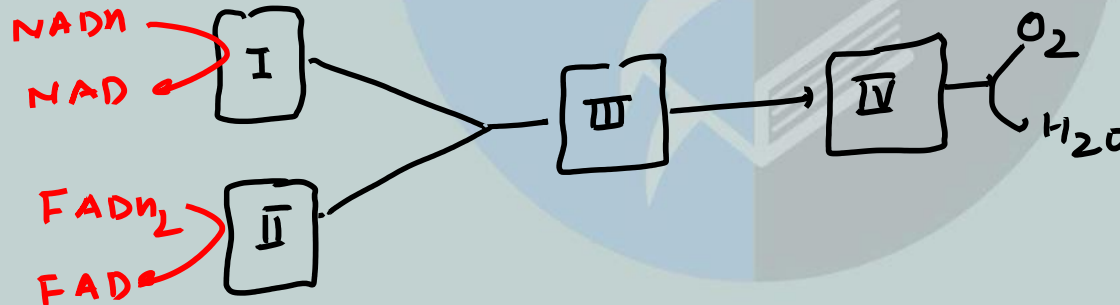
⊥[⊖]
Complex II

Phenformin
Dimercaprol
Antimycin A
BAL (British Anti
Lewisia)

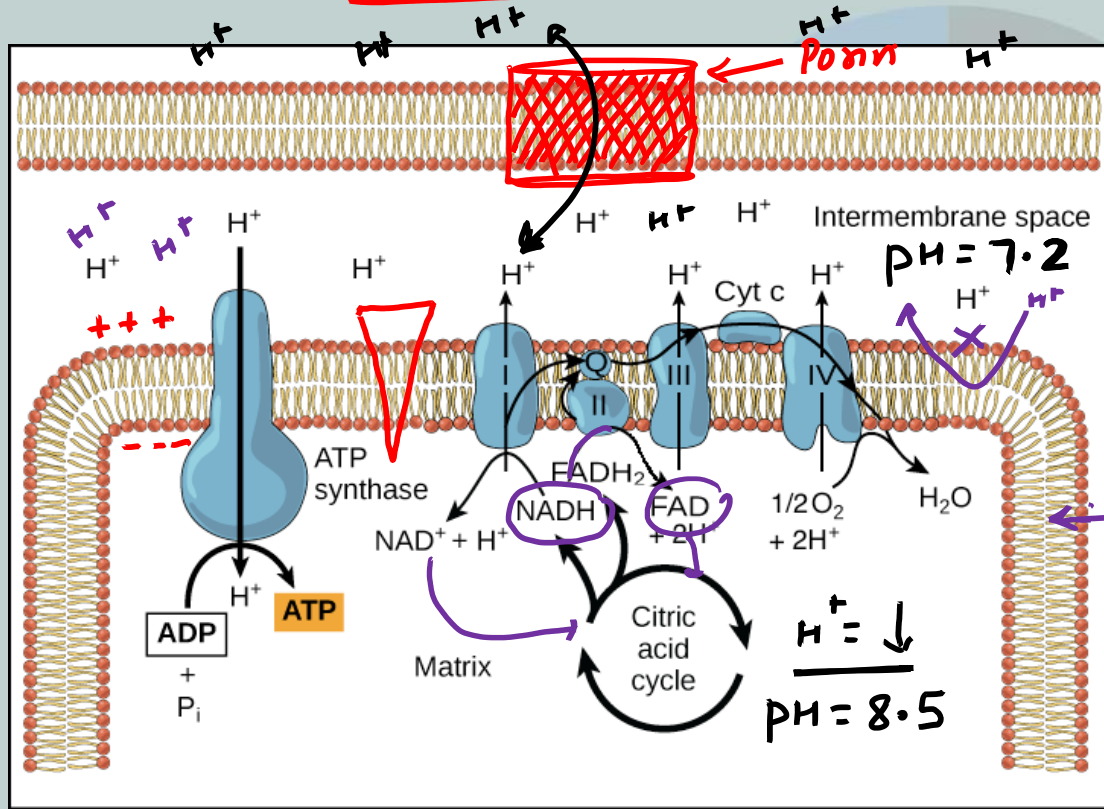
⊥[⊖]
Complex III

Cyanide
Azide
CO
H₂S

⊥[⊖]
Complex-IV



Formation of electrochemical (H^+) gradient



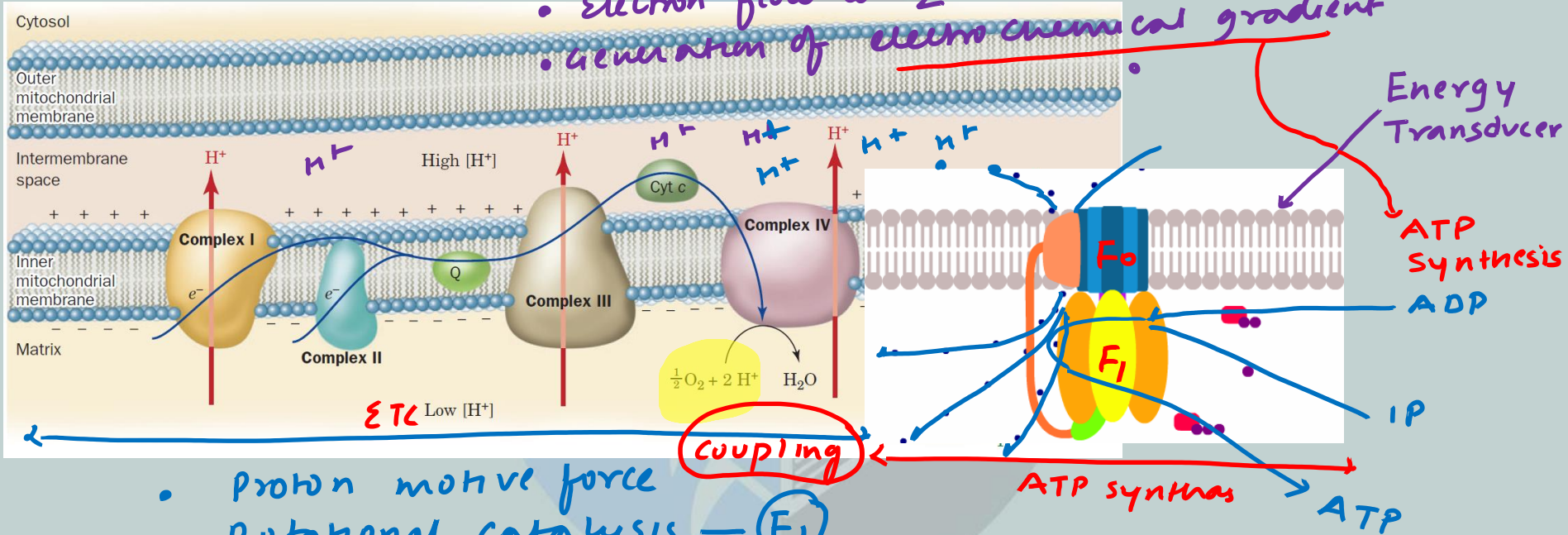
- Cytosol or intermembrane space has around 20 times more H^+ as compared to mitochondrial matrix

Inner membrane is impermeable to H^+



Oxidative phosphorylation:

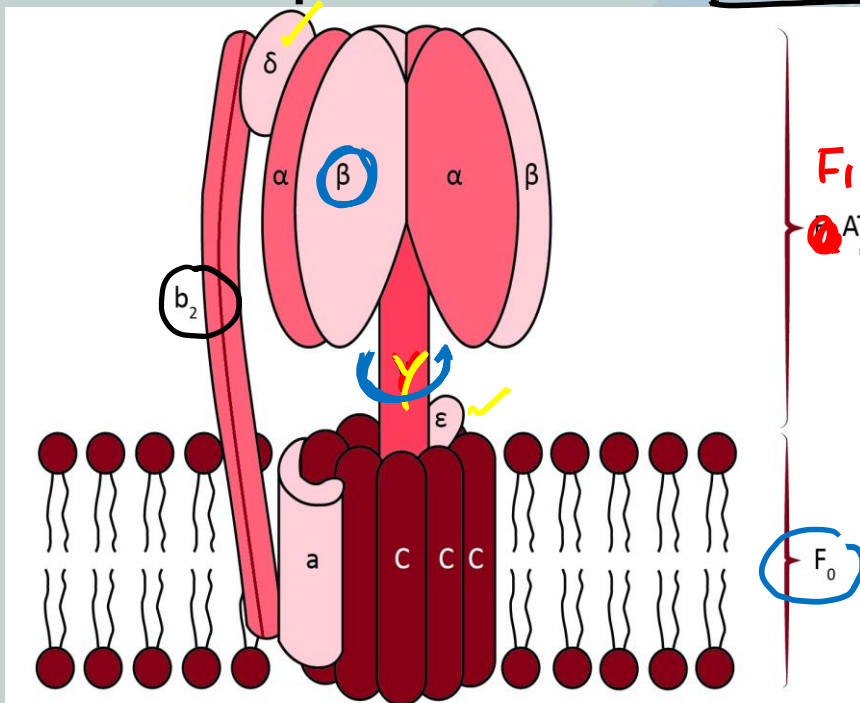
- Oxidation of biomolecule in TCA, β -oxidation
- Production of FADH_2 & NADH
- FADH_2 & NADH are oxidized by complex I or II
- Electron flow to O_2
- Generation of electrochemical gradient



- Proton motive force
- Rotational catalysis — (F_1)
- Phosphorylation of ADP to produce ATP

ATP Synthase Is Driven by the Flow of Protons

The F₁ Component Has Pseudo-Threefold Symmetry
 The F₀ Component Includes a Transmembrane Ring



F₁ ← Peripheral protein
 [3α], [3β] → Catalytic
 1 γ, 1 δ, 1 ε

1 a, 2 b

10 c-subunit

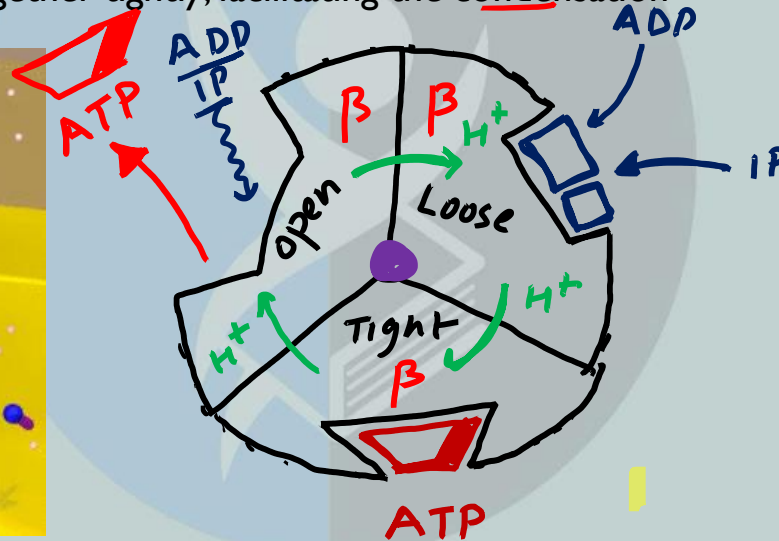
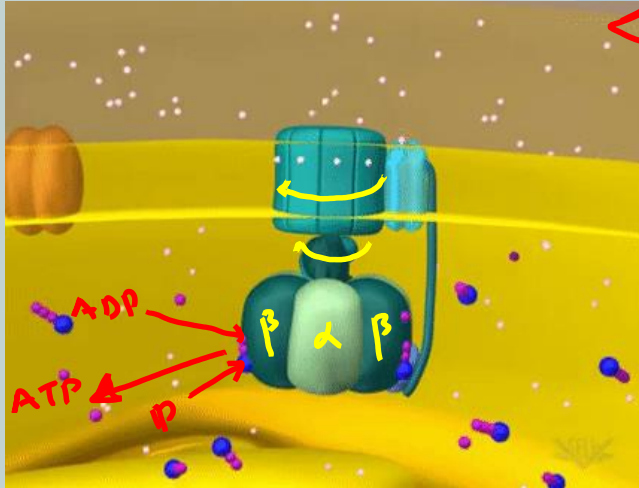
↓
 F₀ channel



ATP Is Synthesized by the Binding Change Mechanism.

Each of the β (nucleotide binding sites) three subunits cycle through 3 different conformations due to rotation of γ subunit:

- ✓ **(O)pen:** little affinity; ADP and P_i are free to enter (ATP is free to leave) the catalytic site
- ✓ **(L)oose:** higher affinity; lose binding of ADP and P_i
- ✓ **(T)ight:** Packing the ADP and P_i together tightly; facilitating the condensation



$3 H^+$ are needed
for 1 ATP
synthesis by
Fo - F₁ particle
(ATP Synthase)
(F-Type of ATPase)

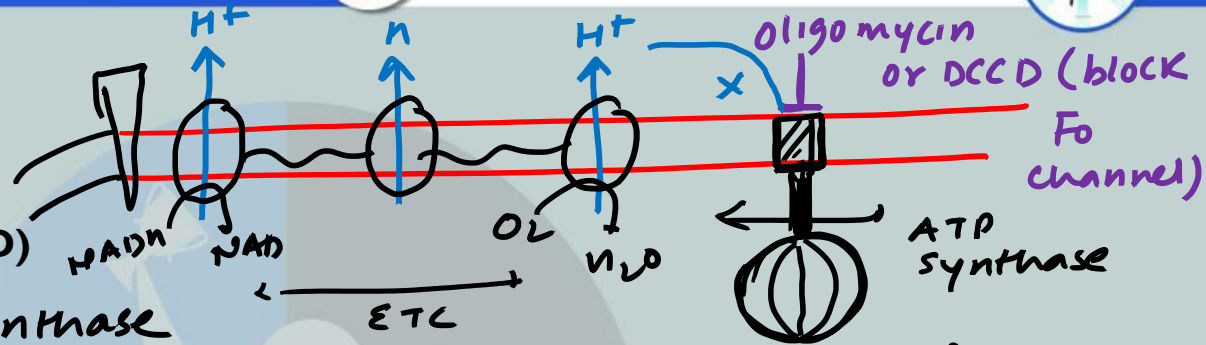


ATP synthase Inhibitors

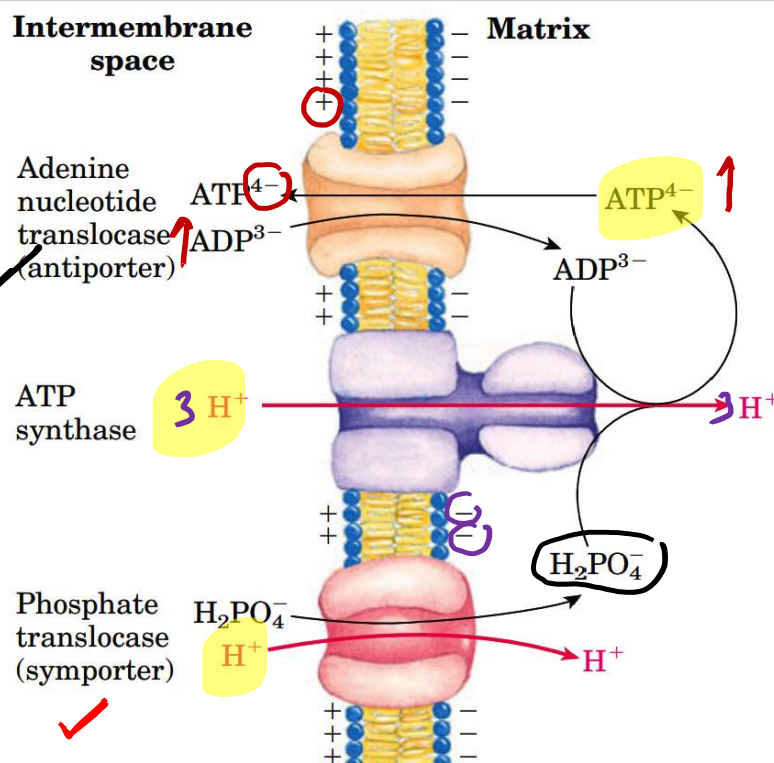
[Oligomycin
Dicyclohexylcarbodiimide (DCCD)]

Directly inhibit ATP synthase

but since electron transport chain is coupled with ATP synthesis thus ETC will also pause.



ADP/ATP transport systems



ADP/ATP Translocator

- 1 ADP (-3) is exchanged for 1 ATP (-4)
- No energy investment
- Along electrochemical gradient movement

Phosphate Translocator

- movement of phosphate is along chemical gradient but opposite to electrical gradient
 - Co-transport : H⁺ / phosphate
- Energy of H⁺ gradient is used for active transport of (P)

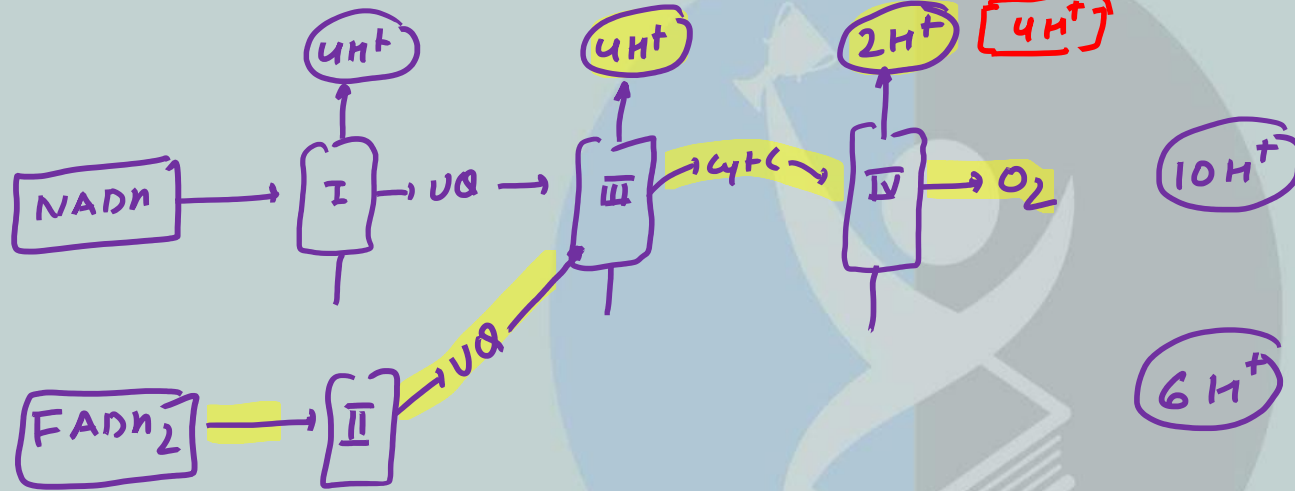


Atractyloside and bongkrekic acid/ bongkrekate inhibit ADP/ATP translocator



The free energy from NADH to O_2 can drive the synthesis of approximately 2.5 ATP

The free energy from $FADH_2$ to O_2 can drive the synthesis of approximately 1.5 ATP

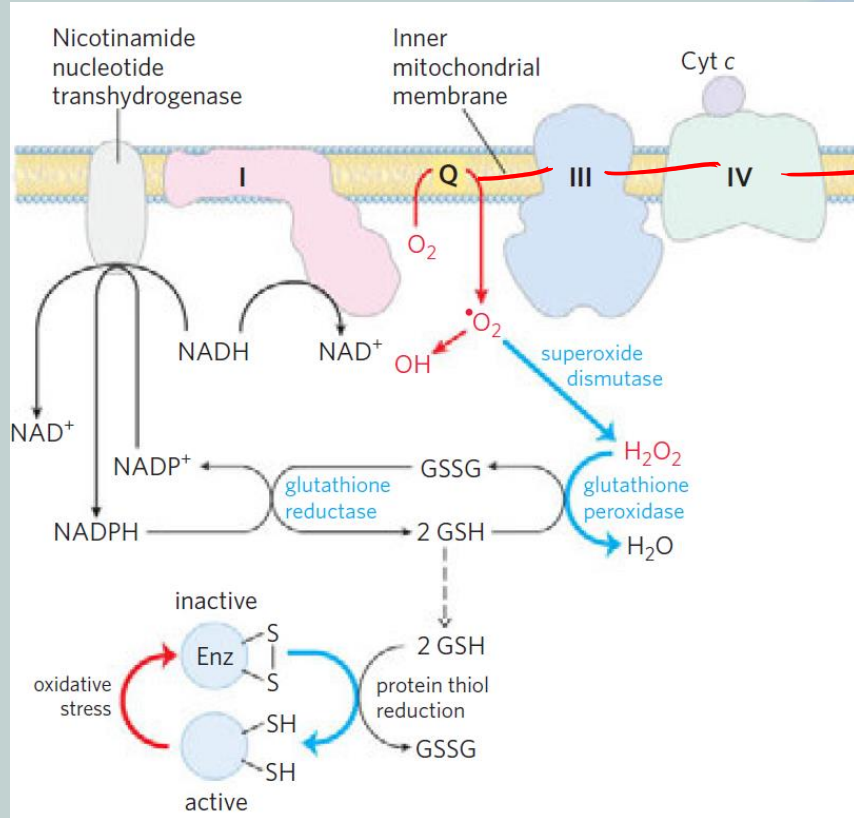


$$4H^+ = 1 \text{ ATP}$$

$$\frac{10H^+}{4H^+} = 2.5 \text{ ATP}$$

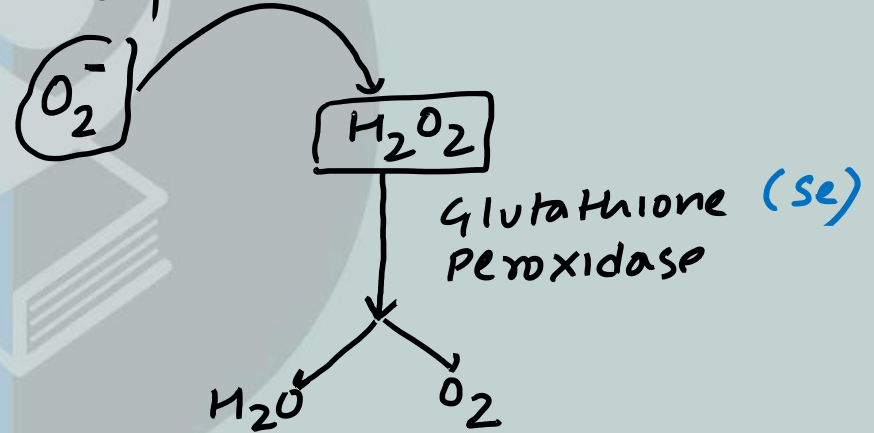
$$\frac{6H^+}{4H^+} = 1.5 \text{ ATP}$$

Reactive Oxygen Species Are Generated during Oxidative Phosphorylation



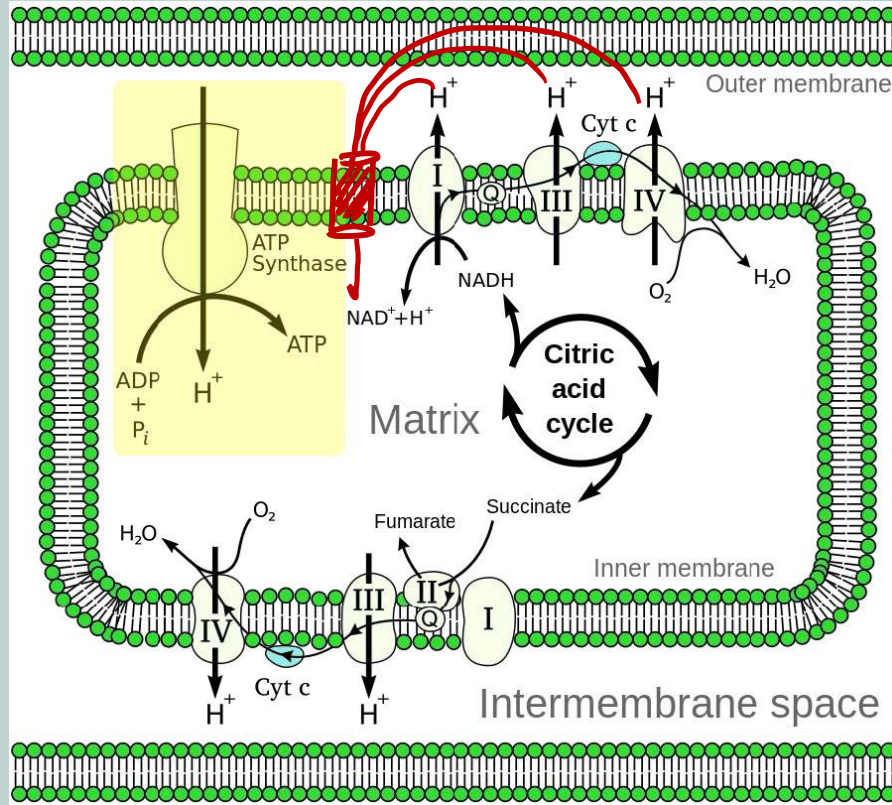
- Anti-oxidant machinery
- Neutralize ROS

Super oxide Dismutase (Mn^{2+})





Coupling of ETC with ATP synthesis:

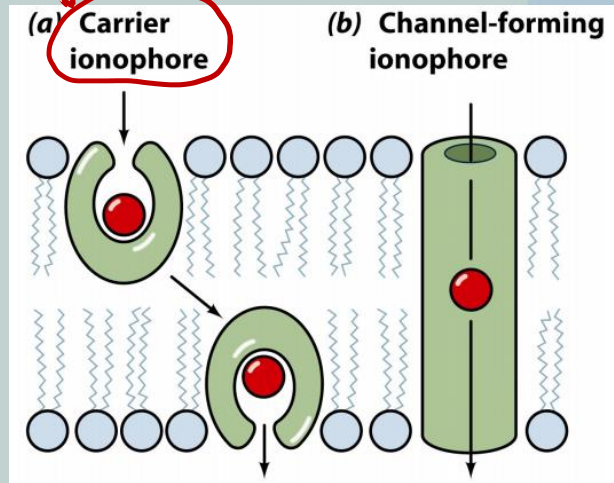


- ETC can function only if H^+ enters back to matrix through F_o channel of ATP synthase.
- Inner memb^r is impermeable to H^+ .

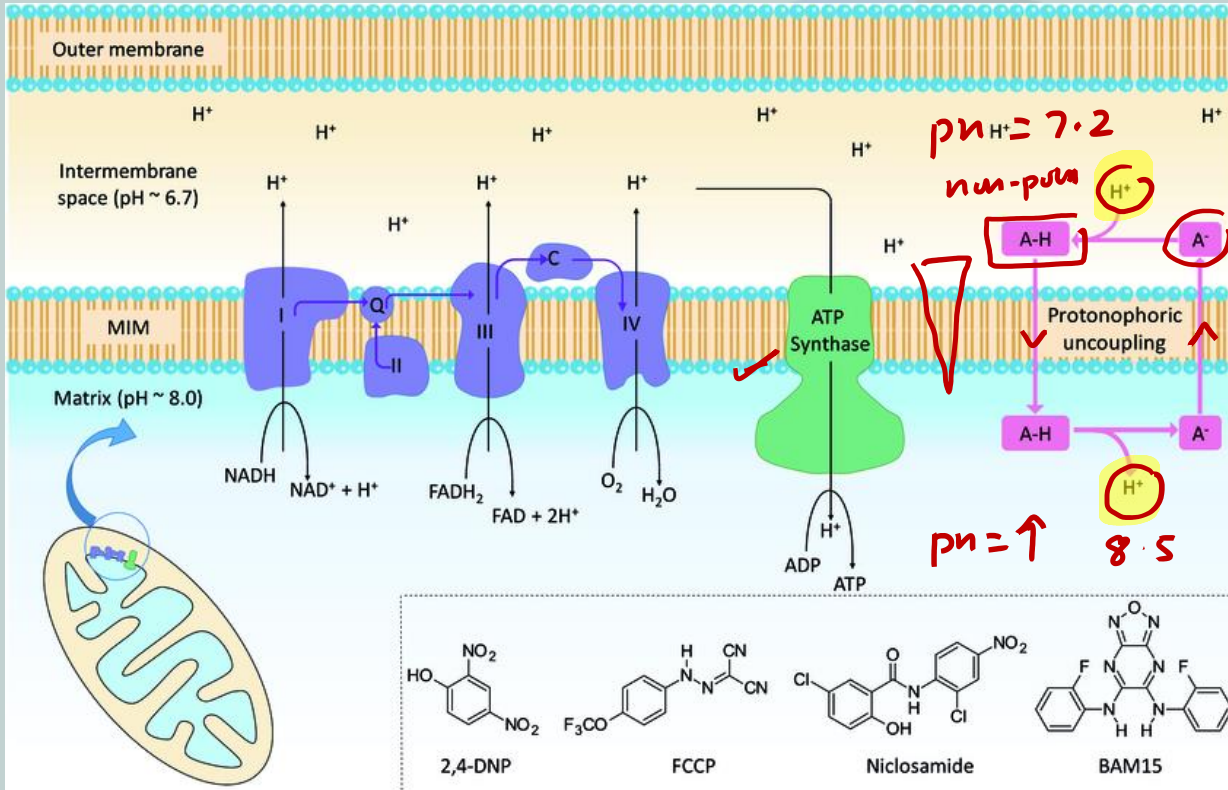
Uncouplers

- Allow electron transport chain to operate in absence of ATP synthesis
- Provide alternative route for entry of H^+ to mitochondrial matrix
- Energy is released in form of heat

1. Ion carrier
2. Ionophore — channel forming



Synthetic uncouplers (Ion carriers):



✓ **2,4-dinitrophenol:**
proton carrier

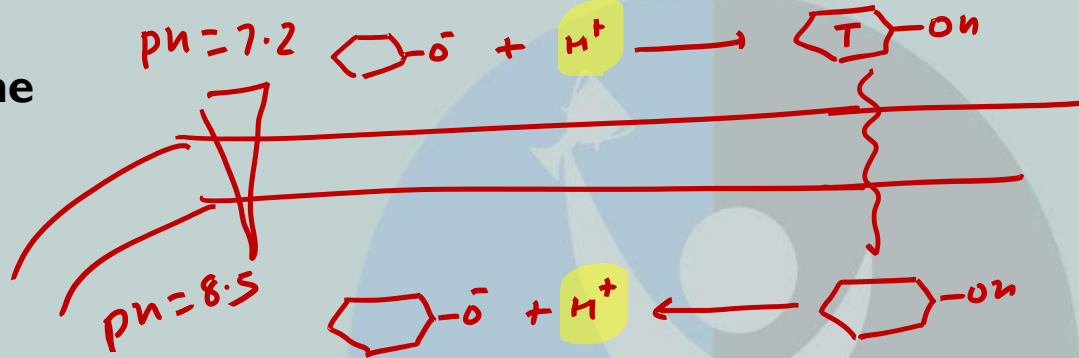
→ FCCP
→ Aspirin
→ Niclosamide



Physiological uncouplers (ion carriers)

Thyroxine

Bilirubin

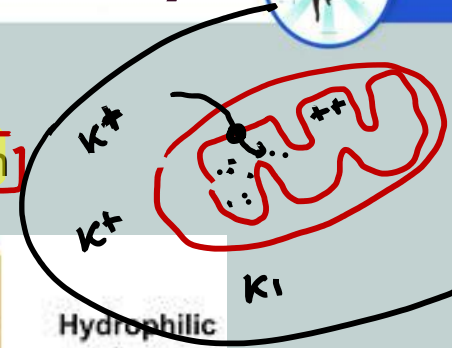
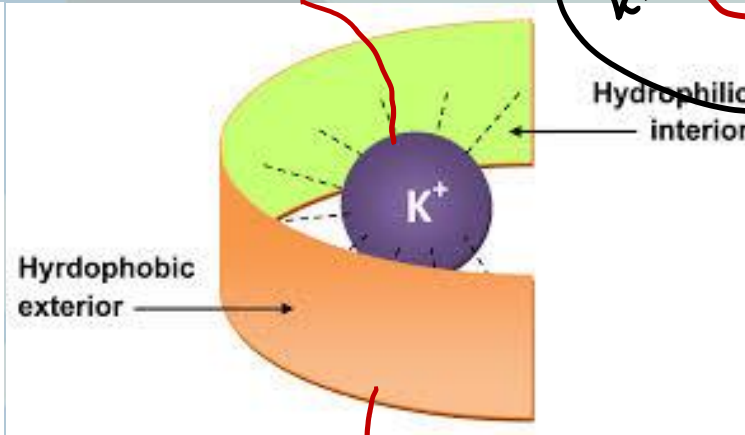
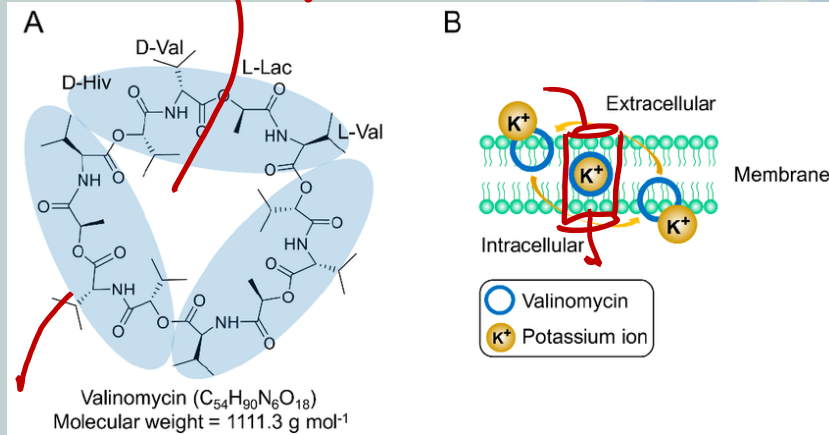


ATP Generation = NO
 Heat Generation = YES
 ETC operate = YES
 O₂ consumption = YES.

Ionophores

Antibiotics **valinomycin** and **nigericin** act as ionophores for **K⁺ ion**

*D & L-amino acid
Amphipathic pore forming helix*



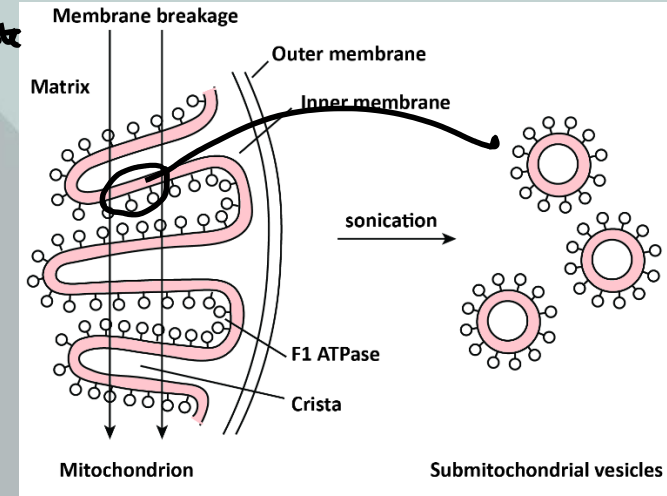
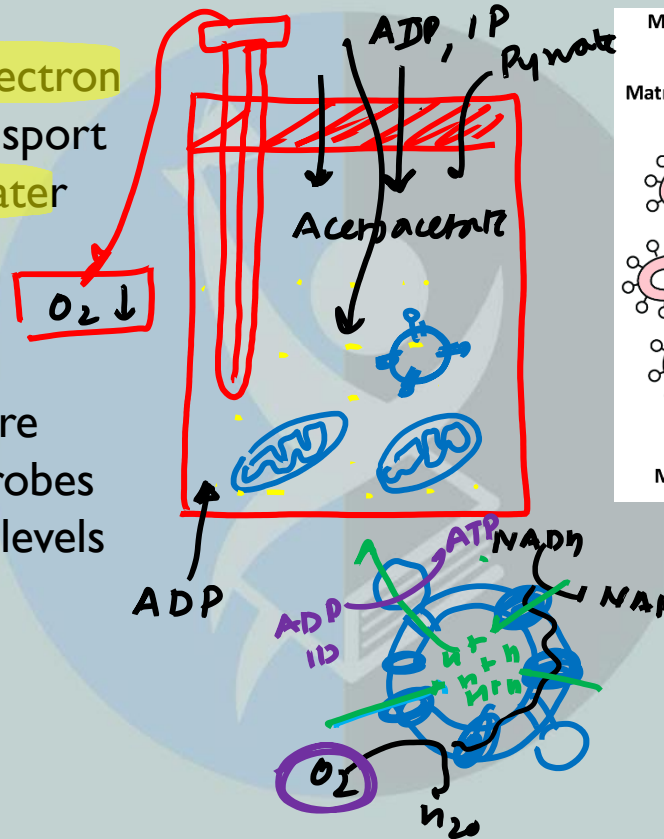
Brown adipose tissue: In polar Animals

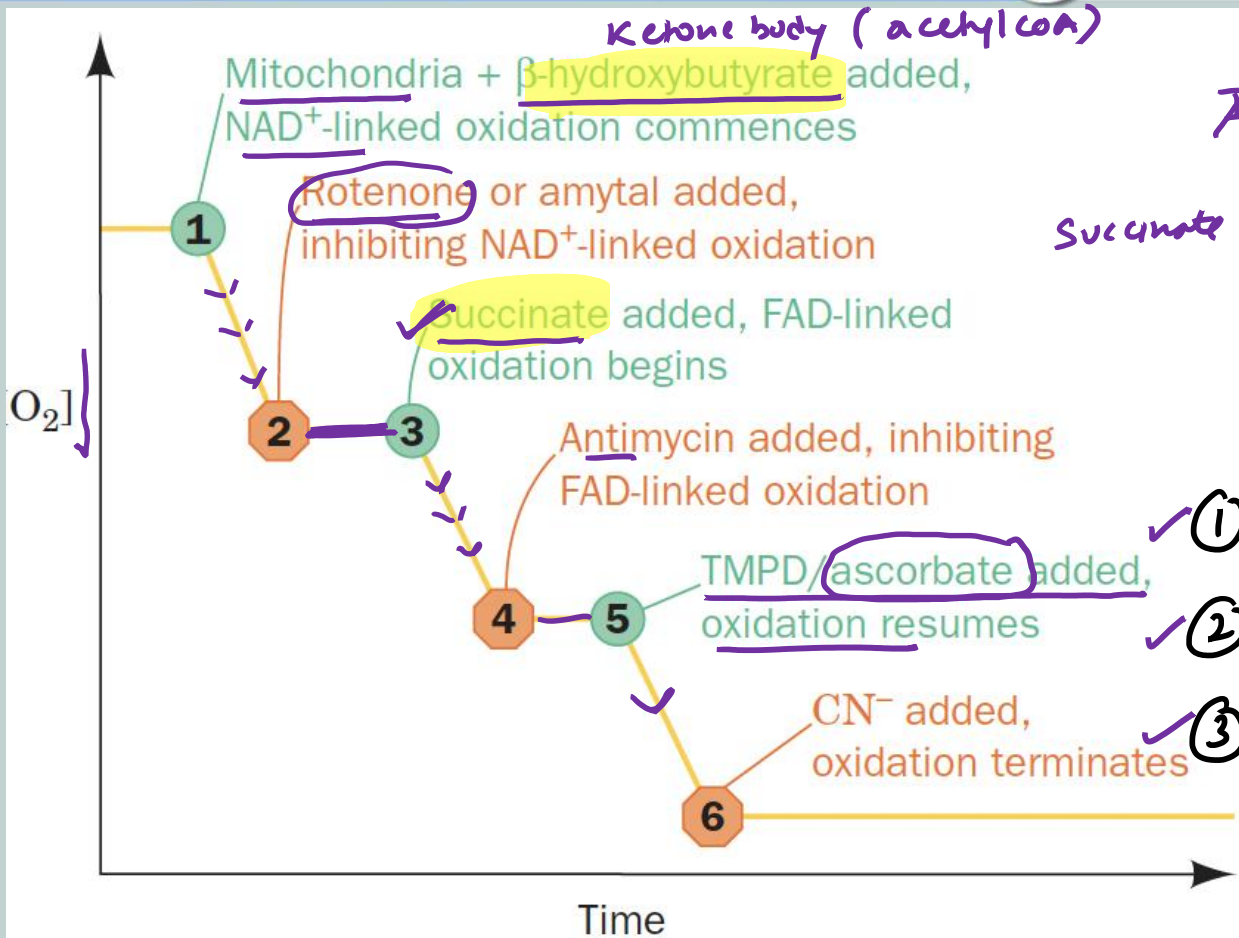
Thermogenin (or uncoupling protein). : *Ionophore for H⁺ ions*

Study of effect of various molecules on electron transport chain

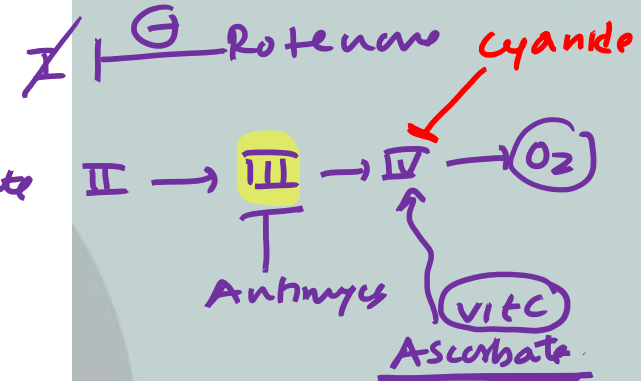
Oxygen serves as the final electron acceptor in the electron transport chain, and its reduction to water marks the completion of the electron transport process.

Clark-type electrodes or more advanced oxygen-sensitive probes are used to measure oxygen levels in a closed system.





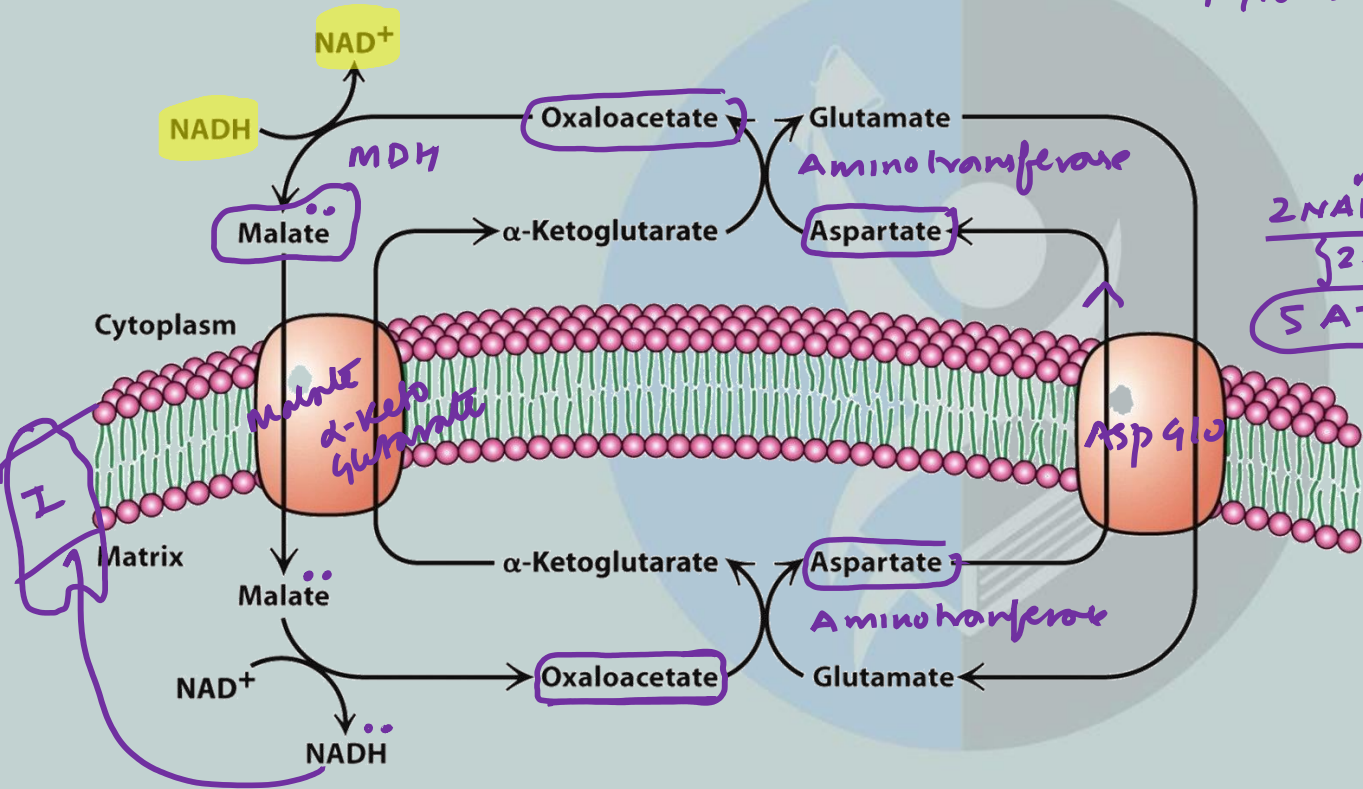
Succinate



Rotenone	off	} inhibit Complex I
Succinate	on	
Antimycin	off	} inhibit complex III
Ascorbic acid	on	
Oligomycin	off	} inhibit ATP Synthase
2,4 DNP	on	



Malate-Aspartate Shuttle (Heart, liver and kidney):



Glycolysis

1 Glucose \rightarrow 2 Pyruvate

2 ATP
2 NADH (cytosol)

2 NADH $\xrightarrow{2.5}$ 5 ATP

ETC mitochondria
Inner memb^v

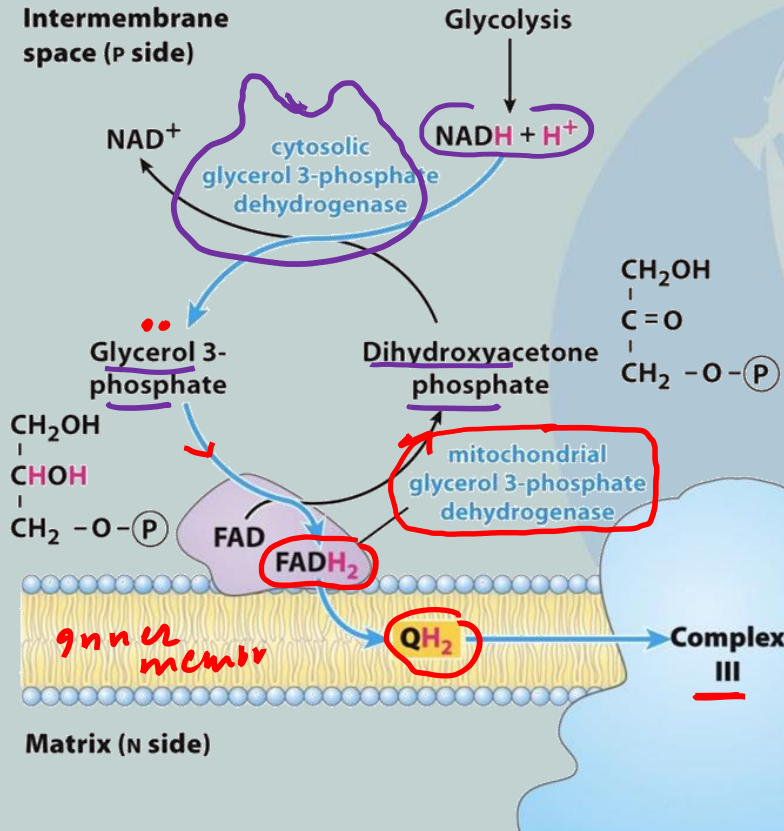
1 Glucose

Glycolysis = 7 ATP

2 Pyruvate = 25 ATP

32 ATP

The glycerophosphate shuttle (brain and muscles); plants



cytosolic
Glycerol-3-P DH \xrightarrow{NADH} NAD^+

mitochondrial
Glycerol-3-P DH \xrightarrow{FAD} $FADH_2$

2 NADH \rightsquigarrow 2 FADH₂
 \downarrow 1.5 ATP
3 ATP

1 glucose

• Glycolysis = 2 + 3 = 5 ATP
• 2 Pyruvate = 25 ATP
30 ATP



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